

636  
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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/939,581A

DATE: 11/13/2001  
TIME: 15:14:26

Input Set : N:\Crf3\RULE60\09939581A.txt  
Output Set: N:\CRF3\11132001\I939581A.raw

4 <110> APPLICANT: Hermeking, Heiko  
5       Vogelstein, Bert  
6       Kinzler, Kenneth  
8 <120> TITLE OF INVENTION: 14-3-3 SIGMA ARREST THE CELL CYCLE  
11 <130> FILE REFERENCE: 1107.77810  
13 <140> CURRENT APPLICATION NUMBER: 09/939,581A  
14 <141> CURRENT FILING DATE: 2001-08-28  
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17 <151> PRIOR FILING DATE: 1998-12-15  
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25 <212> TYPE: DNA  
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34 aggggcccga ggtgcgtgag taccgggaga aggtggagac tgagctccag ggcgtgtgcg     360  
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37 gtgacgacaa gaagcgcatac attgactcag cccgggtcagc ctaccaggag gcoatggaca     540  
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39 ccgtcttcca ctacgagatc gccaacagcc ccgaggaggc catctctctg gccaagacca     660  
40 ctttcgacga ggcatggct gatctgcaca ccctcagcga ggactccatc aaagacagca     720  
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42 aagagggggg cgaggctccc caggagcccc agagctgagt gttggccgccc accgccccgc     840  
43 cctgccccct ccagtcccccc accctgccc gaggactagt atgggggtggg agggccccacc     900  
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62 Gly Glu Glu Leu Ser Cys Glu Glu Arg Asn Leu Leu Ser Val Ala Tyr  
 63 35 40 45  
 64 Lys Asn Val Val Gly Gly Gln Arg Ala Ala Trp Arg Val Leu Ser Ser  
 65 50 55 60  
 66 Ile Glu Gln Lys Ser Asn Glu Glu Gly Ser Glu Glu Lys Gly Pro Glu  
 67 65 70 75 80  
 68 Val Arg Glu Tyr Arg Glu Lys Val Glu Thr Glu Leu Gln Gly Val Cys  
 69 85 90 95  
 70 Asp Thr Val Leu Gly Leu Leu Asp Ser His Leu Ile Lys Glu Ala Gly  
 71 100 105 110  
 72 Asp Ala Glu Ser Arg Val Phe Tyr Leu Lys Met Lys Gly Asp Tyr Tyr  
 73 115 120 125  
 74 Arg Tyr Leu Ala Glu Val Ala Thr Gly Asp Asp Lys Lys Arg Ile Ile  
 75 130 135 140  
 76 Asp Ser Ala Arg Ser Ala Tyr Gln Glu Ala Met Asp Ile Ser Lys Lys  
 77 145 150 155 160  
 78 Glu Met Pro Pro Thr Asn Pro Ile Arg Leu Gly Leu Ala Leu Asn Phe  
 79 165 170 175  
 80 Ser Val Phe His Tyr Glu Ile Ala Asn Ser Pro Glu Glu Ala Ile Ser  
 81 180 185 190  
 82 Leu Ala Lys Thr Thr Phe Asp Glu Ala Met Ala Asp Leu His Thr Leu  
 83 195 200 205  
 84 Ser Glu Asp Ser Tyr Lys Asp Ser Thr Leu Ile Met Gln Leu Leu Arg  
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 99 gacctctttt ctacatagtc ttttttaat ggaaggagaa aatgtcagcc acattactgt 180  
 100 ctgtgttagt ccaggtgaag gtttatcaga aggctgggtg gttttaataa gtttatttcca 240  
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 103 tgagtatcag aagcactctg agccaagggg acagggggca cgtcactgg tcacgagaaa 420  
 104 accctgggtt cccactgggg ctcagccccag cctccttatct ttcccttcc tatggacttc 480  
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115	ctac	cttta	ttaagccag	tattcttgc	tcctgctgt	aataaaactt	1140
116	gagg	ttgtcttt	gtttgggtt	gtttttgtt	tgctttcct	ttgctgaggc	1200
117	agcc	ctgt	tcttcagac	aaatttgggtt	ctttcctggg	gagactgtga	1260
118	agcc	ca	tctggctaca	tttccctca	cctggctgga	gctctgtccg	1320
119	agc	agagagg	gctggggctg	agccccatg	ggcacgtgaa	aagaggccat	1380
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121	act	gccttc	ca	cactccgatc	cccaatga	gaaaatgttt	1500
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123	agcc	agacaa	ctcataaac	tggcccac	ctctggtac	tccctcagga	1620
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126	tct	catc	c	atcg	catg	tcgccaaccc	1800
127	gtgc	gggtgt	tgt	gttgc	gttgc	gttgcagggt	1860
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135	ggg	ctta	agg	ac	ctgc	ccaggaatgg	2340
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145	ccgt	gcac	cc	ccat	ac	ccat	2940
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153	aac	gat	ctc	tgg	atgc	tttgc	3420
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155	cgag	tttcc	cc	tttgc	tttgc	tttgc	3540
156	agat	gagg	gg	actg	gg	tttgc	3600
157	aagc	agca	aa	tcc	gg	tttgc	3660
158	ccca	agg	aaat	gat	gg	tttgc	3720
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